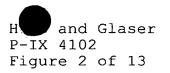


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Figure la

CAG Gln 1	GTG Val	CAG Gln	CTG _. Leu	GTG Val 5	GAG Glu	TCT Ser	GGG Gly	GGA Gly	GGC Gly 10	GTT Val	GTG Val	CAG Gln	CCT Pro	GGA Gly 15	AGG Arg		48
TCC Ser	CTG Leu	aga Arg	CTC Leu 20	TCC Ser	TGT:	GCA \Ala	GCC Ala	TCT Ser 25	GGA Gly	TTC Phe	ACC Thr	TTC Phe	AGT Ser 30	AGC Ser	TAT Tyr		96
GAC Asp	ATG Met	TCT Ser 35	TGG Trp	GTT Val	CGC Arg	CAG Gln	GCT Ala 40	CCG Pro	GGC Gly	AAG Lys	GGT Gly	CTG Leu 45	GAG Glu	TGS TIP	GTC Val	1	.44
SCA Ala	AAA Lys So	GTT Val	AGT Ser	AGT Ser	GGT Gly	GGT Gly 55	GG,Т Gly	AGC Ser	ACC Thr	TAC Tyr	TAT Tyr 60	TTA Leu	GAC Asp	ACT Thr	GTG Val	1	.92
CAG Gln 65	GGC Gly	CGA AEg	TTC Phe	ACC Thr	ATC Ile 70	TCC Ser	AGA Arg	GAC Asp	TAA neA	AGT Ser 75	AAG Lys	OAA neA	ACC Thr	CTA Leu	TAC TYT 08	2	40
CTG Leu	CAA Gln	ATG Het	AAC Asn	TCT Ser 85	CTG Leu	AGA Arg	GCC Ala	GAG Glu	GAC Asp 90	ACA Thr	GCC Ala	GTG Val	TAT TYE	TAC Tyr 95	TGT Cys	· 2	88
GCA Ala	AGA Arg	CAT	AAC Asn 100	TAC Tyr	GGC Gly	AGT Ser	TTT Phe	GCT Ala 105	TAC Tyr	TGG Trp	GGC Gly	CAA Gln	GGG Gly 110	ACT Th:	ACA Thr		36
GTG /al	ACT Thr	GTT Val	TCT Ser	AGT Ser												3	51



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~									Fi	gur	≥ Lb					
				3	•				1.0			. Leu	ser	3:0	GGA Gly	
			20	,	1			25	_			116	26I	Asn	CAC His	95
		,,,					40	-			110	AEG	reu	Leu	ATC Ile	144
						55		-	ATC Ile	~ = 0	60	AL G	rue	ser	Gly	192
					70				ACT Thr	75	002	Set	reu	GTff	Pro	240
				0.0					CAG Gln 90		GGC Gly	AGC Ser	TGG Trp	Pro	CAC His	288
ACG '	TTC Phe	GGA Gly	GGG Gly 100	GCG GGG	ACC Thr	AAG	var	GAA Glu 105	ATT	AAG Lys				95		321

Figure 2a

GA	1 GTG	7 C 3	~ ~~							-						
				•					1 0)		- Ly:	2 5 5 5 6	3 G F	A AGG Y Arg	
			_					25	•				, 261	Se:	TAT	
							40				9	7160	Giu	7, 23	GTC Val	144
						55					- 1 - 2	7191	va b	Tra	GTG Val	192
GLn 65	GLY	CGA Arg	TTC Phe	ACC Thr	ATC Ile 70	TCC Ser	AGA Arg	GAC Asp	AAT Asn	GCC Ala	AAG Lys	AAC Asn	ACC Thr	CTA Leu	TAC Tyr	240
CTG Leu	CAA Gln	ATG Met	AGC Ser	AGT Ser 85	CTG Leu	AAC Asn	TCT Ser	GAG Glu	GAC Asp	ACA Thr	GCC Ala	ATG Met	TAT Tyr	TAC Tyr	TGT Cys	288
GCA .	AGA	Car	3 3 4					GCT Ala 105					GGG GLy	93		336
GTC :	ACT Thr	GTC	ىكى بىل	CC	•			- -					110			351

Figure 2b

				5					1.0		. 561	Val	THE	, bro	GGA Gly	
			20			,		25				116	ser	Yau	CAC His	. 96
		2,2					40				CCA Pro	45	rea	Leu	Ile	144
						55		•			TCC Ser 60	ALG	rne	Ser	GLY	192
					70					75	AAC Asn	Ser	val	GTn.	Thr	240
				0.3					90		GGC Gly	AGC Ser	TGG Trp	CCT Pro 95	CAC His	282
lcg Thr	TTC Phe	gga Gly	GGG Gly 100	GLY	ACC Thr	AAG Lys	CTG Leu	GAA Glu 105	ATT Ile	AAG Lys				<i>3</i> 3		321

LM609 Competition Assay

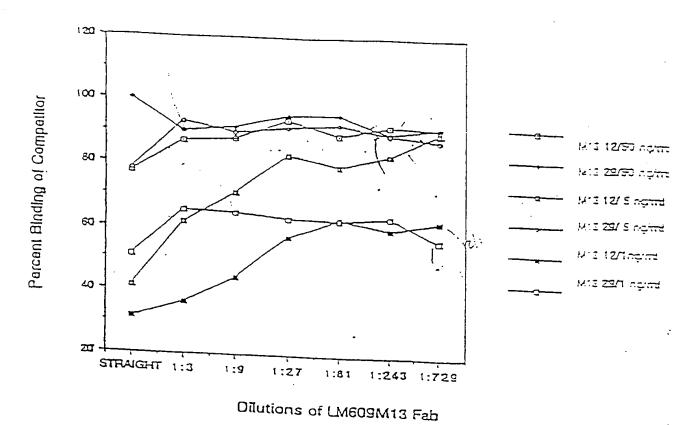
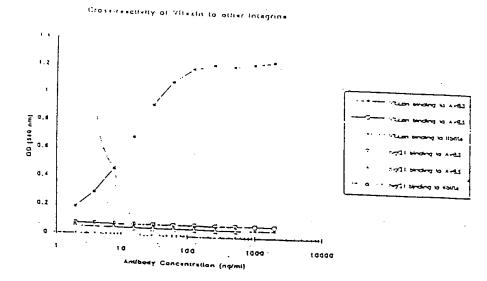


Figure 3

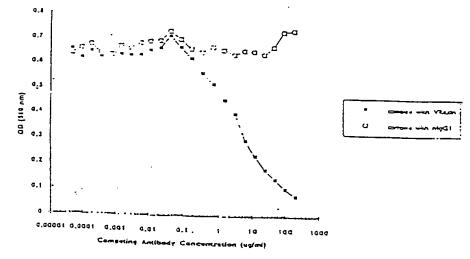


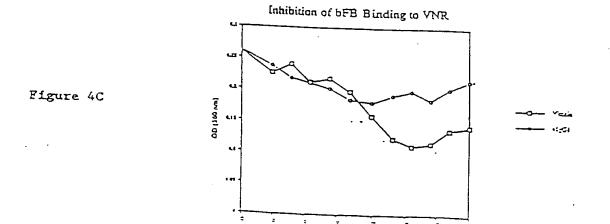
Figure 4A

Pigure 4B



LM609/Vitaxin Competition for Binding to Av83





Inhibition of M21 Melanoma Cell Adhesion to Fibrinogen

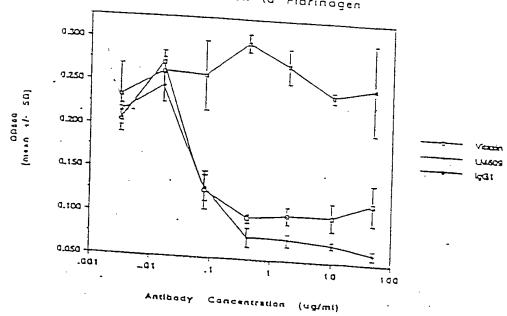
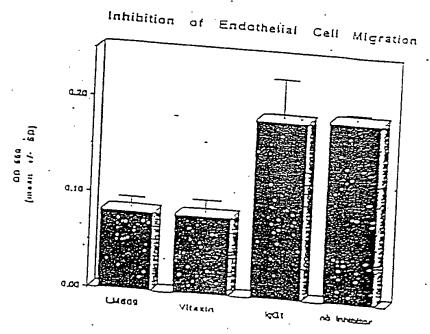


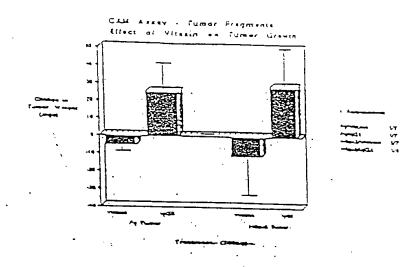
Figure 5B

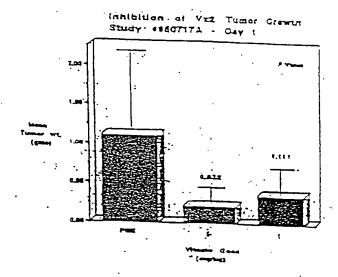
Figure 5A



Treetment







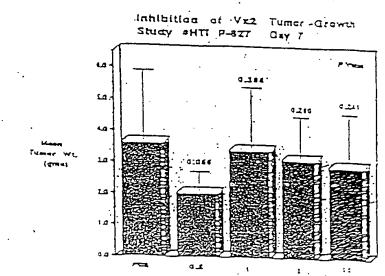


Figure 6B

Figure 6A

Figure 6C

H and Glaser P-IX 4102 Figure 9 of 13

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Figure 7

GAG AT			5	- ÷	301	FLO	Ala	LO	Lou	ser	Lau	Ser	250	GLY		48
GAA AGG Glu Ard	GCG J Ala	ACT Thr 20	CTT Leu	TCC Ser	TGC Cya '	CAG Gln	GCC Ala 25	AGC Ser	CAA Gln	AGT Ser	ATT Ile	AGC Ser 30	AAC Asn	E7a CYC		96
CTA CAC Leu Bis	TGG Trp 35	TAT Tyr	CAA Gln	CAA Gln	AGG AFģ	CCT Pro 40	GGT Gly	CAA Gln	GCC Ala	CCA Pro	AGG Arg 45	CTT Leu	CTC Lau	ATC Ile		144
CGT/ATC Arg/Met	TAT TYP 50	CGT AFG	TCC Ser	CAG Gln	TCC Ser	ATC Ile 55	TCT Ser	GGG Gly	ATC Ile	CCC Pro	GCC Ala 60	AGG Arg	TTC Phe	AGT Ser	GGC Gly	192
AGT GG: Ser Gl; 65	TCA Ser	GGG Gly	ACA Thr	GAT Asp 70	TTC Phe	ACC Thr	CTC Lau	ACT Thr	ATC Ile 75	TCC Ser	AGT Ser	CTG Leu	GAG Glu	30 30		240
GAA GAT	TTT Phe	GCA Ala	GTC Val 85	TAT Tyr	TAC Tyr	TGT Cya	CAA Gln	CAG Gln 90	AGT Ser	GGC Gly	AGC Ser	TGG Trp	CCT Pro 95	CAC Els		288
ACG TTO Thr Phe	GGA Gly	GGG Gly 100	GGG GLY	ACC Thr	AAG Lys	GTG Val	GAA Glu 105	ATT Ile	AAG Lys							321

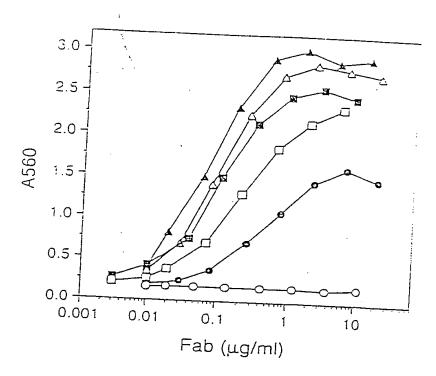


FIGURE 8



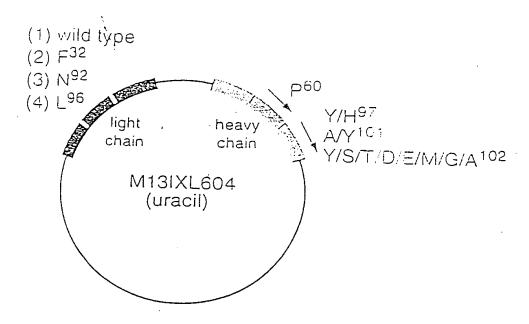
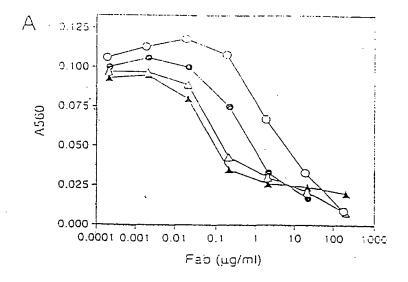


FIGURE 9



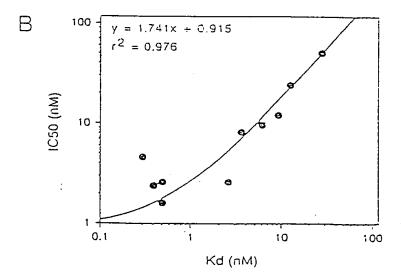


FIGURE 10

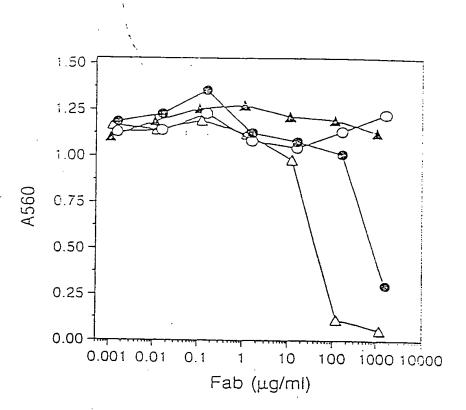


FIGURE 11